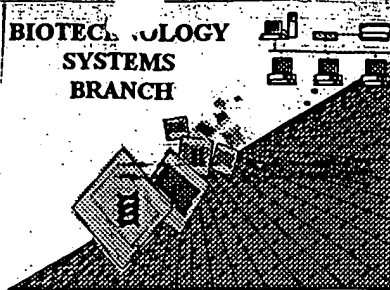


RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/676,380

Source: OIPE

Date Processed by STIC: 10-10-00

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin30help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO).

Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/676,380

DATE: 10/10/2000
TIME: 16:02:47

Input Set : A:\ES.txt
Output Set: N:\CRF3\10102000\I676380.raw

**Does Not Comply
Corrected Diskette Needed**

2 <110> APPLICANT: Maihle, Nita Reiter, Jill Baron, Andre
4 <120> TITLE OF INVENTION: SOLUBLE EPIDERMAL GROWTH FACTOR RECEPTOR-LIKE PROTEINS AND THEIR USES IN
5 CANCER DETECTION METHODS
7 <130> FILE REFERENCE: Atairgin 257/225
C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/676,380
C--> 9 <141> CURRENT FILING DATE: 2000-09-29
9 <150> PRIOR APPLICATION NUMBER: 60/157,144
W--> 11 <151> PRIOR FILING DATE: 09/30/99
E--> 13 <160> NUMBER OF SEQ ID NOS: 20 <170> PatentIn version 3.0<210> 1<211> 705<212> PRT<213> homo

↑ ↑ ↑ ↑ ↑
ERRORED SEQUENCES

*Enter "hard return" at these
positions.*

✓ F.Y.I.

**The types of errors shown exist throughout the Sequence Listing. Please check
subsequent sequences for similar errors.**

VERIFICATION SUMMARY

DATE: 10/10/2000

PATENT APPLICATION: US/09/676,380

TIME: 16:02:48

Input Set : A:\ES.txt

Output Set: N:\CRF3\10102000\I676380.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:11 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:14 M:282 W: Numeric Field Identifier Missing, <210> is required.
L:14 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:14 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:14 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:14 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQ ID NO
L:14 M:283 W: Missing Blank Line separator, <400> field identifier
L:13 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (20) Counted (1)